Title: 47324, A NOVEL HUMAN G-PROTEIN AND USES THEREFOR

Inventors: Rachel Meyers

Application No.: To be Assigned

Docket No.: 381552003500

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					M	A	S	L	D	R	V	K	v	L	v	L	G	D	s	SEQ ID NO:	2
AGG	GAAG	GCAG	rggc.	AAG	AIG (الرك ا	TCC	CTG	GAT	CGG	GTG	AAG	GTA (CTG	GTG	TTG	GGA	GAC	TCA	SEQID NO:	i
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GGT	GTT	GGG	AAA	TCT	TCG	TTA.	GTC	CAT	CTC	CTA	TGC	CAA	AAT	CAA	GTG	CTG	GGA	AAT	CCA	105	
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GAA	GAG	AAG	ACC	TAC	TAC	ATA	GAA	TTA	TGG	GAT	GTT	GGA	GGC	TOT	GTG	CCC	≥ C.T.	CCC	AGC	225	
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AGC	GTG	AAA	AGC	ACA	AGA	GCA	GTA	TTC	TAC	AAC	TCC	GTA	AAT	GGT	ATT	ATT	TTC	GTA	CAC	285	
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	L	T	N	K	K	S	5	Q	N	L	R	R	W	S	L	E	A	L	N	115	
GAL	TTA	AĻA	AAT	AAG	AAG	TCC	100	CAA	AAC	TTG	CGT	CGT	TGG	TCA	TTG	GAA	GCT	CTC	AAC	345	
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AGG	GAT	TTG	GTG										GAT	TAT	GAT	240	C2 2	CYC	TTT	405	
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GCT	GAT	AAC	CAA	ATA	CCA	CTG	TTG	GTA	ATA	GGG	ACT	AAA	CTG	GAC	CAG	ATT	CAT	GAA	ACA	465	
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D	*																			237	
GAC	TGA																			711	
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47324	2	*->GlgflkiflkllglwnKEmRILiLGLDnAGKTTILykLklge.ivtt SEQ ID NO:4 ++1 + +++L+LG + GK+ +++ L +++ + + ASLDRVKVLVLGDSGVGKSSLVHLLCQNQvLGNP 35
47324	36	<pre>iPTiGFNvEtveykNikFtvWDvGGQeslRPlW + T+G v v + + ++++++++ + WDvGG ++ ++ s R + SWTVGCSV-DVRVHDykegtpeektYYIELWDVGGSvgsassvKSTRAV- 83</pre>
47324	84	<pre>rnYfpnTdavIfVVDSaDreRieeakeELhaLLneeeL + +++++IfV D ++ + + L + e +++ +++FYNSVNGIIFVHDLTNKKSSQNLRRWSLEALNrdlvptgvlvtn 127</pre>
47324	128	adAplLifANKQDlpgAmseaEireaLgLhelkgskg ++ ++++ + plL+ + K D + +E+ +++ L + gdydqeqfadNQIPLLVIGTKLDQIHETKRHEVltrTAFLAEDFNP-E 174
47324	175	kVtl.e.gdRpWeiqgcsAvkGeGLyEGldWLsnnikkr<-* ++ l+ ++ R + ++Avk l + +++++ EINLdCtnPRYLAAGSSNAVKLSRFFDKVIEK 206

FIG. 3

47324	8	*->KlVligDsgVGKSsLliRFtdnkFveeYipTIGvDFytktvevD SEQ ID NO:5 K++++GDsGVGKSsL++ +n+ T+G+ ++ ++ ++ KVLVLGDSGVGKSSLVHLLCQNQVLGNPSWTVGCSVDVRVHDYKegt 54
47324	55	GktvkLqIWDTAGQERFralrpaYYRgAqGfLLVYDITs ++ kt+ ++WD G ++ +a++ ++++ +Y +G++ V D T+ peEKTYYIELWDVGGSVGSAssvkstravFYNSVNGIIFVHDLTN 99
47324	100	rdSFenvkkWleeilrhadkdenvpivLVGNKc ++S +n+++W e l+++ +++ ++++ ++++ d+ +p + +G+K KKSSQNLRRWSLEALNrdLVPtgvlvtngdydqeqfADNQIPLLVIGTKL 149
47324	150	DLeddedlelte.gqkRvVsteeGealAkelgAlpFmETSA D + e++ +V t ++ lA++++ ++ ++ +++ + DQIHEtKR-HEVLTRTAF-LAEDFNPeeinldctnPRYLAAGS 190
47324	191	KtntNVeeaFeelareilkkvsevnvnldqpakkkkskCcil<-* +++ F+ ++++ + ++++++ + +k+ + SNAVKLSRFFDKVIEKRYFLREGNQIPGFPD-RKRFGAGTLK 231

FIG. 4

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Query: Sbjct: NO:6	35 PSWTV-GCSVDVRVHDYKEGTPEEKTYYIELWDXXXXXXXXXXXXXTRAVFYNSVNGI 91 P+WT S D + Y TP Y++E +D R FY +++GI 69 PTWTTPSSSEDSENYPYMRSTPTTTNILYFVEFYDLNSDWRMCRQQRESFYKNIDGI 125 SEQ ID
Query:	92 IFVHDLTNKKSSQNLRRWSLEALNRDLVPTGVLVTNGDYDQEQFADNQIPLLVIGTKLDQ 151 + V+++ S +L W + L + + + + + P+LV+GT LD+
Sbjct:	126 VLVYNMLELSSQDSLHDWLYDPLRQICKHRHLRIRSILKNHNAPILVVGTNLDK 179
Query:	152 IHETKRHEVLTRTAFLAEDFNPEEINLDCTNPRYLAAGSSNAVKLSRFFDKVIE-KRYF 209 + L R +A N EE+ ++C +P+ S N KL F ++VIE K F
Sbjct:	180 LMRRPLRRRGSIAHQLNVEEMLVNCLDPQSFVDKSRNQGKLYGFLNRVIEFKEQF 234

FIG. 5a

Ouery:	7 VKVLVLGDSGVGKSSLVHLLCQNQVLGNP-SWTVGCSVDVRVHDYKE 52
21	V++L+LGD GVGK+SL +L+ ++ P S TVG V VR+H+Y +
Shict:	12 VRILMLGDRGVGKTSLTNLMATTEITPTPDSRTVGEESWHVQVRLHEYSK 61 SEQ ID NO:

FIG. 5b

Query:		TGVLVTNGDYDQEQFAD-NQIPLLVIGTKLDQIHETKRHEVLTRTAFLAEDFNPEE 17: T L T+G +D E+F Q P+LV+GTKLD + E KRH + + +A+ EE		
Sbjct: No:8	19	TDNLGTDGHILFDMEEFLGATQTPILVMGTKLDLLDE-KRHPKMGVKKPGGIADKCGAEE 77	SEQ	ID
Query:	176	INLDCTNPRYLAAGSSNAVKLSRFFDKVIEKRYFLREGNQIPGFPDRKRFGAG 22 I L+C N R LAAG+++AVKLSRFFD+VIE R LR + PDR+RFG	3	
Sbjct:	78	IWLNCRNSRSLAAGTTDAVKLSRFFDRVIENRKALRAALAFGVSSSNAVSPPDRRRFGPT 13	7	
Query:	229	TLK 231 + K		
Sbjct:	138	SAK 140		

FIG. 6

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6 RVKVLVLGDSGVGKSSLVHLLCQNQVLGNPSWTVGCSVDVRVHDYKEGTPEEKTYYIE 63 Query:

+V+VLV+GDSGVGK+SLVHL+ + + P T+GC+V V+ Y G+P + I+

80 QVRVLVVGDSGVGKTSLVHLINKGSSIVRPPQTIGCTVGVKHITY--GSPASSSSSIQ 135 SEQ ID NO:9 Sbjct:

FIG. 7

8 KVLVLGDSGVGKSSLVHLL--CQNQ------VLGNPSWTVGCSVDVRVHDYKEGT 54 Query: V+ + T+G +V+ K++++GDSGVGK+SL++ L +N 7 KIVIIGDSGVGKTSLLNKLRFTENSFTEEYDPTTRTVVDSYKSTIGVDFNVKTTIEVVDT 66 SEQ ID Sbjct: NO:10 55 PEEKTYYIELWDXXXXXXXXXXXXTRAVFYNSVNGIIFVHDLTNKKSS-QNLRRWSLEA 113 Query: I V+D+ + +SS +N +W E A + Y+ K +++WD 67 -DGKNIKLQIWDTAGQERYRSMLTSMEA-YYRGAEADIIVYDVDSSESSFENQTKWLKEI 124 Sbjct: 114 LNRDLVPTGVLVTNGDYDQEQFADNQIPLLVIGTKLD-QI---HETKRHEVLTRTAFLAE 169 Query: +N E+ ++N +P++++G K D ++ E + + T 125 LRH-----ASN----EEASEN-VPIILVGNKADLEVPNPEEVEEEKEEASTEEEAQ 170 Sbjct:

170 DFNPEEINLDCTNPRYLAAGSSNAVKLSRFFDKVIEKRYFLREGNQIPGFPDRKRF 225 Query:

F +++ R L++ +I P + F EE L 171 SF-AEEKGLGVV-PFIETSAKTTGTNVEEVFQELV--REILKKKKEIQEKADQEKY 222 Sbjct:

FIG. 8